

Figure 1

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Figure 2A

Figure 2B

		10	20	30	40	50	
HUMAN	1	MLLSKFGSLA	HLCGPGGVDH	LPVKILOPAK	ADKESFEKAY		50
RAT	1	MLLSKFGSLA	HLCGPGGVDH	LPVKILOPAK	ADKESFEKVY	OVGAVLGSGG	50
MOUSE	1	MLLSKFGSLA	HLCGPGGVDH	LPVKILOPAK	ADKESFEKVY	OVGAVIGSGG	50
				~		(· u · · · · · · · · · · · · · · · · · · ·	-
		60	70	80	90	100	
HUMAN	51	FGTVYAGSRI	ADGLPVAVKH	VVKERVTEWG	SLGGATVPLE	VVLLRKVGAA	100
RAT	51	FGTVYAGSRI	ADGLPVAVKH	VVKERVTEWG	SLGGMAVPLE	VVLLRKVGAA	100
MOUSE	51	FGTVYAGSRI	ADGLPVAVKH	VVKERVTEWG	SLGGVAVPLE	VVLLRKVGAA	100
		110	120	130	140	150	
HUMAN	101	GGARGVIRLL	DWFERPDGFL	LVLERPEPAQ	DLFDFTTERG	ALDEPLARRF	150
RAT	101	GGARGVIRLL				ALDEPLARRE	150
MOUSE	101	GGARGVIRLL	DWFERPDGFL	LVLERPEPAQ	DLFDFTTERG	ALDEPLARRF	150
		160	170	180	190	200	
HUMAN	151	FAQVLAAVRH				GSGALLKDTV	200
RAT	151	FAQVLAAVRH				GSGAVLKDTV	200
MOUSE	151	FAQVLAAVRH	CHNCGVVHRD	IKDENLLVDL	RSGELKLIDF	GSGAVLKDTV	200
		210	220	230	240	250	
HUMAN	201	YTDFDGTRVY			LGVLLYDMVC		250
RAT	201	YTDFDGTRVY		YHGRSATVWS	LGVLLYDMVC	GDIPFEQDEE	250
MOUSE	201	YTDFDGTRVY	SPPEWIRYHR	YHGRSATVWS	LGVLLYDMVC	GDIPFEQDEE	250
		260	270	280	290	300	
HUMAN	251			RWCLSLRPSE	RPSLDQIAAH	PWMLGADGGA	300
RAT	251	ILRGRLFFRR		EWCLSLRPSE	RPSLDQIAAH	PWMLGTEGSV	300
MOUSE	251	ILRGRLFFRR	RVSPECQQLI	EWCLSLRPSE	RPSLDQIAAH	PWMLGTEGSV	300
		310	320				
HUMAN	301		LDPDDVASTT				
RAT	301	PENCDLRLCA					
MOUSE	301	PENCDLRLCA	LDTDDGASTT	SSSESL			

Figure 3

		10	20	30	40	
Pim-1	1	MLLSKINSLA			40	50
Pim-2	•	MLTKPLQ-	CDD AD	DOTTOTTODOCC	-EFFETTESOA	QVGPLLGSGG 50
Pim-3	;	MLLSKFGSLA		I MOVII ODAY	ADEBUTEANT	RLGPLLGKGG 50
PHD-3		MIASKIGSLA	HIXGINGGVDH	LPVKILQPAK	ADKESFEKAY	QVGAVLGSGG 50
		60	70	80	90	100
Pim-1	51	FGSVYSGIRV				EVVLLKKVSS 100
Pim-2	51		TDRLQVAIKV			EVALLWKVGA 100
Pino-3						EVVLLREVGA 100
гш-э	31	POLVINGSEL	ALCILLANATU	A AVDVA I PAC	SL-GGAI VPL	BVVLLKAVGA 100
		110	120	130	140	150
Pim-1	101	GFSGVIRL				GALQEELARS 150
Pim-2	101		LDWFRTORGE	MI.VI.ERPI.PA	ODITEDALLER	GPLGEGPSRC 150
Pim-3	101	AGGARGVIRL	LDWFERPDGF	LLVLERPEPA	ODLEDENER	GALDEPLARR 150
					dom or time	CHADILITIES 150
		160	170	180	190	200
Pim-1	151	FFWOVLEAVR	HCHNCGVLHR			FGSGALLKDT 200
Pim-2	151		HCHSRGVVHR			FGSGALLHDB 200
Pim-3	151		HCHSCGVVHR			FGSGALLKDT 200
		•				
		210	220	230	240	250
Pim-1	201	VYTDFDGTRV	YSPPEWIRYH	RYHGRSAAVW	SLGILLYDMV	CGDIPFEHDE 250
Pim-2	201	PYTDFDGTRV	YSPPEWISRH	QYHALPATVW	SLGILLYDMV	CGDIPFERDO 250
Pim-3	201	VYTDFDGTRV	YSPPEWIRYH	RYHGRSATVW	SLGVLLYDMV	CGDIPFEODB 250
					•	•
		260	270	280	290	300
Pim-1	251	BURGQVFFR	QRVSSECQHL	IRWCLALRPS	DRPTFEEIQN	HPWMQDVLL- 300
Pim-2	251	BILRARLHFP				DPWMQTPABD 300
Pim-3	251	BILRGRLLFR	RRVSPECQQL	IRWCLSLRPS	ERPSLDQIAA	HPWMLGADGG 300
		310	320	330	340	
Pim-1		PQET	ARIHL	HSLSPG	P	SK
Pim-2	301	VTPQPLQRRP	CPFGLVLATL			
Pim-3	301	APBS	CDLRL	CTLDPD	DVASTTSSSE	SL

Figure 4

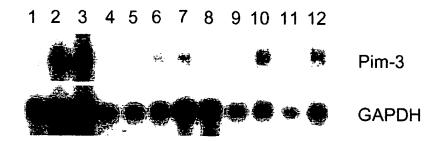
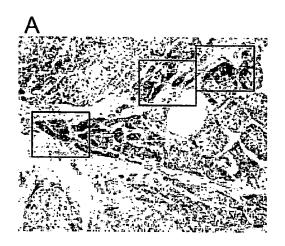


Figure 5



Figure 6



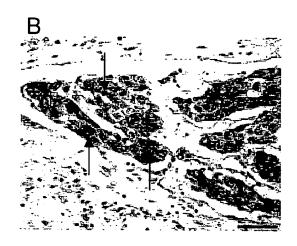


Figure 7

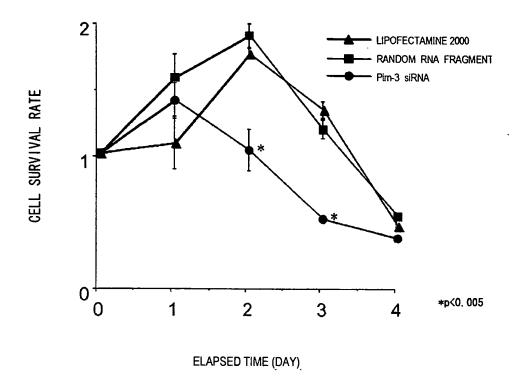


Figure 8

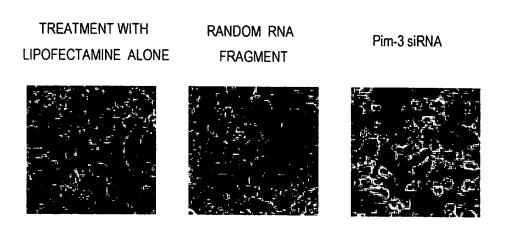


Figure 9

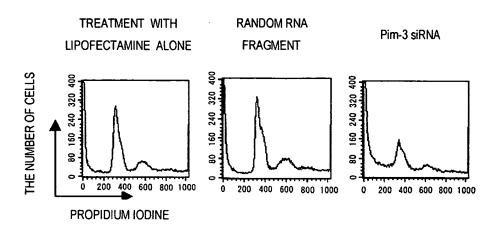


Figure 10